

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:	10/514,626
Source:	PCT
Date Processed by STIC:	08/04/2005
	7 7 7

ENTERED



PCT

RAW SEQUENCE LISTINGPATENT APPLICATION: **US/10/514,626**DATE: 08/04/2005

TIME: 14:41:08

Input Set : A:\sequence listing.txt
Output Set: N:\CRF4\08042005\J514626.raw

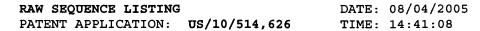
4 <110> APPLICANT: Pipeline-Biotech

```
Finn Skou Pedersen
     6
             Shervin Bahrami
     7
             Mogens Ryttergaard Duch
     9 <120> TITLE OF INVENTION: A purified retroviral envelope
     10
             polypeptide, isolated nucleic acids encoding said
     11
             polypeptide, vectors and use thereof
    13 <130> FILE REFERENCE: P32118PC01
C--> 15 <140> CURRENT APPLICATION NUMBER: US/10/514,626
C--> 15 <141> CURRENT FILING DATE: 2004-11-17
     15 <150> PRIOR APPLICATION NUMBER: PA200200767
     16 <151> PRIOR FILING DATE: 2002-05-17
     18 <160> NUMBER OF SEQ ID NOS: 2
     20 <170> SOFTWARE: FastSEQ for Windows Version 4.0
     22 <210> SEQ ID NO: 1
    23 <211> LENGTH: 1920
    24 <212> TYPE: DNA
    25 <213> ORGANISM: Murine Leukaemia Virus
    27 <400> SEOUENCE: 1
    28 atggaaggtc cagcgttctc aaaacccctt aaagataaga ttaacccgtg gggcccccta 60
     29 atagteetgg gaatettaat gagggeaaga gtateagtae aacatgaeag eeeteateag 120
     30 gtcttcaatg ttacttggag agttaccaac ttaatgacag gacaaacagc taatgctacc 180
     31 teceteetgg ggacaatgae egatgeettt eetaaaetgt aetttgaett gtgegattta 240
    32 ataggggacg actgggatga gactggactc gggtgtcgca ctcccggggg aagaaaaagg 300
    33 gcaagaatat ttgacttcta tgtttgcccc ggtcacactg tgctagcagg gtgtggaggg 360
    34 ccgagagagg gctactgtgg caaatgggga tgtgagacca ctggacaggc atactggaag 420
    35 ccatcatcat catgggacct aattteeett aagegaggaa acaeteetaa aggeeaggge 480
    36 ccctgttatg attecteggt ggtetecagt agegeecagg gtgecacace ggggggtega 540
     37 tgcaaccccc tagtcctaga attcactgac gcgggtaaaa gggccagctg ggacgcctcc 600
    38 aaagcatggg gactaagact gtaccgatcc acaaggaccg acccggtgac ccggttctct 660
    39 ttgacccgcc aggtcctcaa tatagggccc cgcgtcccca ttgggcctaa tcccgtgatc 720
    40 attgaccagt tacccccctc ccgacccgtg cagatcatgc tccccaggcc tcctcagcct 780
    41 cctccaccag gcgcagcctc tacagtccct gagactgccc caccttccca acaacctggg 840
    42 acgggagaca ggctgctaaa cctggtaaat ggagcctacc aagctctcaa cctcaccagt 900
    43 cctgacaaaa cccaagagtg ctggttgtgt ctggtagcgg gaccccccta ctacgaaggg 960
    44 gttgccgtcc taggtactta ttccaaccat acctctgccc cagctaactg ctccgtggcc 1020
    45 teccaacaca agetgaceet gteegaagtg aceggacagg gaetetgegt aggageagtt 1080
    46 cccaaaaccc atcaggccct gtgtaatacc acccagaaga cgagcaacgg gtcctactat 1140
    47 ctggctgctc ccgccgggac catttgggct tgcaacaccq qqctcactcc ctgcctatct 1200
    48 accactgtgc tcgacctcac caccgattac tgtgtcctgg ttgagctctg gccaaaagtg 1260
    49 acctaccact cccctggtta tgtttatggc cagtttgaag aaaaaaccaa atataaaaga 1320
    50 gaaccegtet cactaactet ggeeetacta ttaggaggae teactatggg eggaattgee 1380
    51 geoggagtgg gaacagggac tacegeecta gtggeeacte ageagtteea acaacteeag 1440
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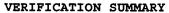
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52 gctgccatgc aggatgacct taaagaagtt gaaaagtcca tcactaatct agaaagatct 1500 53 ttgacctcct tgtccgaagt agtgttacag aatcgtagag gcctagatct actattccta 1560 54 aaagagggag gtttgtgtgc tgccttaaaa gaagaatgct gtttctatgc cgaccacaca 1620 55 ggattggtac gggatagcat ggccaaactt agagaaagat tgagtcagag acaaaaactc 1680 56 tttgaatccc aacaagggtg gtttgaaggg ctgtttaaca agtccccttg gttcaccacc 1740 57 ctgatatcca ccatcatggg tcccctgata atcctcttgt taattttact ctttgggcct 1800 58 tgtattetea ateaeetggt eeagtttate aaagaeaggg ttteggtagt geaggeeetg 1860 61 gtcctgactc aacaatatca tcaacttaag acaatagaag attgtgaatc acgtgaataa 1920 63 <210> SEQ ID NO: 2 64 <211> LENGTH: 639 65 <212> TYPE: PRT 66 <213> ORGANISM: Murine Leukaemia Virus 68 <400> SEQUENCE: 2 69 Met Glu Gly Pro Ala Phe Ser Lys Pro Leu Lys Asp Lys Ile Asn Pro 10 71 Trp Gly Pro Leu Ile Val Leu Gly Ile Leu Met Arg Ala Arg Val Ser 73 Val Gln His Asp Ser Pro His Gln Val Phe Asn Val Thr Trp Arg Val 35 40 75 Thr Asn Leu Met Thr Gly Gln Thr Ala Asn Ala Thr Ser Leu Leu Gly 55 77 Thr Met Thr Asp Ala Phe Pro Lys Leu Tyr Phe Asp Leu Cys Asp Leu 70 79 Ile Gly Asp Asp Trp Asp Glu Thr Gly Leu Gly Cys Arg Thr Pro Gly 81 Gly Arg Lys Arg Ala Arg Ile Phe Asp Phe Tyr Val Cys Pro Gly His 100 105 83 Thr Val Leu Ala Gly Cys Gly Gly Pro Arg Glu Gly Tyr Cys Gly Lys 120 85 Trp Gly Cys Glu Thr Thr Gly Gln Ala Tyr Trp Lys Pro Ser Ser Ser 135 87 Trp Asp Leu Ile Ser Leu Lys Arg Gly Asn Thr Pro Lys Gly Gln Gly 150 155 89 Pro Cys Tyr Asp Ser Ser Val Val Ser Ser Ser Ala Gln Gly Ala Thr 165 170 91 Pro Gly Gly Arg Cys Asn Pro Leu Val Leu Glu Phe Thr Asp Ala Gly 185 93 Lys Arg Ala Ser Trp Asp Ala Ser Lys Ala Trp Gly Leu Arg Leu Tyr 200 95 Arg Ser Thr Arg Thr Asp Pro Val Thr Arg Phe Ser Leu Thr Arg Gln 215 220 97 Val Leu Asn Ile Gly Pro Arg Val Pro Ile Gly Pro Asn Pro Val Ile 230 235 99 Ile Asp Gln Leu Pro Pro Ser Arg Pro Val Gln Ile Met Leu Pro Arg 245 250 101 Pro Pro Gln Pro Pro Pro Gly Ala Ala Ser Thr Val Pro Glu Thr 265 103 Ala Pro Pro Ser Gln Gln Pro Gly Thr Gly Asp Arg Leu Leu Asn Leu 104 275 280 285



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105 106	Val	Asn 290	Gly	Ala	Tyr	Gln	Ala 295	Leu	Asn	Leu	Thr	Ser 300	Pro	Asp	Lys	Thr
	Gln 305	Glu	Cys	Trp	Leu	Cys 310	Leu	Val	Ala	Gly	Pro 315	Pro	Tyr	Tyr	Glu	Gly 320
		Ala	Val	Leu	Gly 325	Thr	Tyr	Ser	Asn	His 330		Ser	Ala	Pro	Ala 335	
	Cys	Ser	Val	Ala		Gln	His	Lys	Leu		Leu	Ser	Glu	Val	Thr	Gly
112	-			340				-	345					350		-
113	Gln	Gly	Leu	Cys	Val	Gly	Ala	Val	Pro	Lys	Thr	His	Gln	Ala	Leu	Cys
114			355					360					365			
115	Asn	Thr	Thr	Gln	Lys	Thr		Asn	Gly	Ser	Tyr	_	Leu	Ala	Ala	Pro
116		370					375					380	,			
		Gly	Thr	Ile	Trp		Cys	Asn	Thr	Gly		Thr	Pro	Cys	Leu	
	385	1		_	_	390		_,	_	_	395		_			400
	Thr	Thr	Val	Leu		Leu	Thr	Thr	Asp		Cys	Val	Leu	Val	Glu	Leu
123	[]	Desa	T	77-7	405	(Tle	TT: -		D	410		77-3	П	a 1	415	Dha
	ттр	PIO	ьуѕ	420	THE	TYL	HIS	ser	425	GIA	Tyr	vaı	Tyr	430	Gln	Pne
125	Glu	Glu.	Laze		Tarc	Тъгъ	Tvc	7~~		Dro	17a l	Car	T 011		Leu	ח ד ת
127	GIU	Giu	435	1111	цуз	TYL	пуъ	440	Giu	PIO	vaı	SET	445	1111	пеп	AIA
	Len	Len		Glv	Glv	T.e.i	Thr	٠,	Glv	Glv	Tle	Δla		Glv	Val	Glv
129		450		- 1	01		455		U _1	017		460			• • • •	OI,
130	Thr	Gly	Thr	Thr	Ala	Leu	Val	Ala	Thr	Gln	Gln	Phe	Gln	Gln	Leu	Gln
	465	-				470					475					480
132	Ala	Ala	Met	Gln	Asp	Asp	Leu	Lys	Glu	Val	Glu	Lys	Ser	Ile	Thr	Asn
133					485					490					495	
134	Leu	Glu	Arg	Ser	Leu	Thr	Ser	Leu	Ser	Glu	Val	Val	Leu	Gln	Asn	Arg
135				500					505					510		
	Arg	Gly		Asp	Leu	Leu	Phe		Lys	Glu	Gly	Gly		Cys	Ala	Ala
137	_	_	515		_	_		520		_			525	_	-	_
	Leu	_	GIu	Glu	Cys	Cys		Tyr	Ala	Asp	His		GIY	Leu	Val	Arg
139	7	530	Mah	77-	T	T	535	03	7	T	0	540	3	~1	T	Τ
	545	ser	Met	Ala	пĀг	ьеи 550	Arg	GIU	Arg	ьeu	555	GIII	Arg	GIII	Lys	
		Gl 11	Cor	Gln.	G3 n		Trn	Dho	Glu	Glaz		Dho	Nan	Lvc	Ser	560 (
143	FIIC	GIU	261	GIII	565	Gry	тър	FIIC	Giu	570	шец	FIIC	ASII	цуъ	575	PIO
	Trn	Phe	Thr	Thr		Tle	Ser	Thr	Tle	-	Glv	Pro	Len	Tle	Ile	Leu
145				580					585		01,			590		200
	Leu	Leu	Ile		Leu	Phe	Glv	Pro		Ile	Leu	Asn	His		Val	Gln
147			595				2	600	- 2				605			
148	Phe	Ile	Lys	Asp	Arg	Val	Ser	Val	Val	Gln	Ala	Leu		Leu	Thr	Gln
149		610	-	_	_		615					620				
150	Gln	Tyr	His	Gln	Leu	Lys	Thr	Ile	Glu	Asp	Cys	Glu	Ser	Arg	Glu	
151	625					630					635					



PATENT APPLICATION: US/10/514,626

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Input Set : A:\sequence listing.txt
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L:15 M:270 C: Current Application Number differs, Replaced Current Application No

L:15 M:271 C: Current Filing Date differs, Replaced Current Filing Date